

Shibuya

CRF Errors Corrected by the STIC System Branch

1635 #11

Serial Number: 08/886,313

CRF Processing Date: 2/10/2000  
Edited by: AS  
Verified by: \_\_\_\_\_ (STIC staff)

ENTERED

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☒ Other: deleted duplicated seq 5

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

PAGE: 1

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/886,313DATE: 02/11/2000  
TIME: 00:27:56

INPUT SET: S34701.raw

This Raw Listing contains the General  
Information Section and those Sequences  
containing ERRORS.

## SEQUENCE LISTING

Does Not Comply  
Corrected Diskette Needed

--> ok

1  
2  
3 (1) General Information:  
4  
5 (i) APPLICANT: Wright, Jim A.  
6 Young, Aiping H.  
7  
8 (ii) TITLE OF INVENTION: INHIBITING NEOPLASTIC CELLS UNTILIZING  
9 THE RIBONUCLEOTIDE REDUCTASE UTR  
10  
11 (iii) NUMBER OF SEQUENCES: 50  
12  
13 (iv) CORRESPONDENCE ADDRESS:  
14 (A) ADDRESSEE: Kohn & Associates  
15 (B) STREET: 30500 Northwestern Hwy.  
16 (C) CITY: Farmington Hills  
17 (D) STATE: Michigan  
18 (E) COUNTRY: US  
19 (F) ZIP: 48334  
20  
21 (v) COMPUTER READABLE FORM:  
22 (A) MEDIUM TYPE: Floppy disk  
23 (B) COMPUTER: IBM PC compatible  
24 (C) OPERATING SYSTEM: PC-DOS/MS-DOS  
25 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30  
26  
27 (vi) CURRENT APPLICATION DATA:  
28 (A) APPLICATION NUMBER:  
29 (B) FILING DATE:  
30 (C) CLASSIFICATION:  
31  
32 (viii) ATTORNEY/AGENT INFORMATION:  
33 (A) NAME: Kohn, Kenneth I.  
34 (B) REGISTRATION NUMBER: 30,955  
35 (C) REFERENCE/DOCKET NUMBER: 0227.00003  
36  
37 (ix) TELECOMMUNICATION INFORMATION:  
38 (A) TELEPHONE: (248) 539-5050  
39 (B) TELEFAX: (248) 539-5055  
40  
41

ERRORED SEQUENCES FOLLOW:

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/886,313DATE: 02/11/2000  
TIME: 00:27:56

INPUT SET: S34701.raw

--&gt; 894 (2) INFORMATION FOR SEQ ID NO:5:

895

896

(i) SEQUENCE CHARACTERISTICS:

897

(A) LENGTH: 21 base pairs

898

(B) TYPE: nucleic acid

899

(C) STRANDEDNESS: single

900

(D) TOPOLOGY: linear

901

902

903

904

905

906

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

907

908

TAAGTAACTG ATCGTGTGCT C

21

909

*Delete -  
duplicated*

--&gt; 910 (2) INFORMATION FOR SEQ ID NO:50:

911

912

(i) SEQUENCE CHARACTERISTICS:

913

(A) LENGTH: 21 base pairs

914

(B) TYPE: nucleic acid

915

(C) STRANDEDNESS: single

916

(D) TOPOLOGY: linear

917

918

919

920

921

922

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:50:

923

924

TTAAGACTTT TTACGCGATT C

21

925

*OK*

PAGE: 1

**SEQUENCE VERIFICATION REPORT**  
**PATENT APPLICATION US/08/886,313**

DATE: 02/11/2000  
TIME: 00:27:56

**INPUT SET: S34701.raw**

Line	Error	Original Text
11	Number of Sequences (50) Doesn't Equal Actual Count (51)	(iii) NUMBER OF SEQUENCES: 50
894	Wrong Sequence Number	(2) INFORMATION FOR SEQ ID NO:5:
910	Sequence 6 thru 49 missing	(2) INFORMATION FOR SEQ ID NO:50: